SEQUENCE LISTING

<110> Duzic, Emir et al.

<120> AGS PROTEINS AND NUCLEIC ACID MOLECULES AND USES THEREOF

<130> CPI-086CPPC

<140>
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<160> 72

<170> PatentIn Ver. 2.0

<210> 1
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<213> Homo sapiens

<220>
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Met Lys Leu Ala Ala Met Ile Lys Lys Met Cys Pro Ser Asp Ser Glu
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ctg agt atc ccg gcc aag aac tgc tat cgc atg gtc atc ctc ggc tcg   96
Leu Ser Ile Pro Ala Lys Asn Cys Tyr Arg Met Val Ile Leu Gly Ser
            20                  25                  30

tcc aag gtg ggc aag acg gcc atc gtg tcg cgc ttc ctc acc ggc cgc  144
Ser Lys Val Gly Lys Thr Ala Ile Val Ser Arg Phe Leu Thr Gly Arg
        35                  40                  45

ttc gag gac gcc tac acg cct acc atc gag gac ttc cac cgc aag ttc  192
Phe Glu Asp Ala Tyr Thr Pro Thr Ile Glu Asp Phe His Arg Lys Phe
    50                  55                  60

tac tcc atc cgc ggc gag gtc tac cag ctc gac atc ctc gac acg tcc  240
Tyr Ser Ile Arg Gly Glu Val Tyr Gln Leu Asp Ile Leu Asp Thr Ser
65                  70                  75                  80

ggc aac cac ccg ttc ccc gcc atg cgg cgc ctc tcc atc ctc aca gga  288
Gly Asn His Pro Phe Pro Ala Met Arg Arg Leu Ser Ile Leu Thr Gly
                85                  90                  95

gac gtt ttc atc ctg gtg ttc agt ctg gac aac cgc gac tcc ttc gag  336
Asp Val Phe Ile Leu Val Phe Ser Leu Asp Asn Arg Asp Ser Phe Glu
            100                 105                 110

gag gtg cag cgg ctc agg cag cag atc ctc gac acc aag tct tgc ctc  384
Glu Val Gln Arg Leu Arg Gln Gln Ile Leu Asp Thr Lys Ser Cys Leu
        115                 120                 125

aag aac aaa acc aag gag aac gtg gac gtg ccc ctg gtc atc tgc ggc  432
Lys Asn Lys Thr Lys Glu Asn Val Asp Val Pro Leu Val Ile Cys Gly
    130                 135                 140

aac aag ggt gac cgc gac ttc tac cgc gag gtg gac cag cgc gag atc  480
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Asn Lys Gly Asp Arg Asp Phe Tyr Arg Glu Val Asp Gln Arg Glu Ile
145                 150                 155                 160

gag cag ctg gtg ggc gac gac ccc cag cgc tgc gcc tac ttc gag atc   528
Glu Gln Leu Val Gly Asp Asp Pro Gln Arg Cys Ala Tyr Phe Glu Ile
            165                 170                 175

tcg gcc aag aag aac agc agc ctg gac cag atg ttc cgc gcg ctc ttc   576
Ser Ala Lys Lys Asn Ser Ser Leu Asp Gln Met Phe Arg Ala Leu Phe
        180                 185                 190

gcc atg gcc aag ctg ccc agc gag atg agc cca gac ctg cac cgc aag   624
Ala Met Ala Lys Leu Pro Ser Glu Met Ser Pro Asp Leu His Arg Lys
    195                 200                 205

gtc tcg gtg cag tac tgc gac gtg ctg cac aag aag gcg ctg cgg aac   672
Val Ser Val Gln Tyr Cys Asp Val Leu His Lys Lys Ala Leu Arg Asn
210                 215                 220

aag aag ctg ctg cgg gcc ggc agc ggc ggc ggc ggc ggc gac ccg ggc   720
Lys Lys Leu Leu Arg Ala Gly Ser Gly Gly Gly Gly Gly Asp Pro Gly
225                 230                 235                 240

gac gcc ttt ggc atc gtg gca ccc ttc gcg cgc cgg ccc agc gta cac   768
Asp Ala Phe Gly Ile Val Ala Pro Phe Ala Arg Arg Pro Ser Val His
                245                 250                 255

agc gac ctc atg tac atc cgc gag aag gcc agc gcc ggc agc cag gcc   816
Ser Asp Leu Met Tyr Ile Arg Glu Lys Ala Ser Ala Gly Ser Gln Ala
            260                 265                 270

aag gac aag gag cgc tgc gtc atc agc tag                          846
Lys Asp Lys Glu Arg Cys Val Ile Ser
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Met Lys Leu Ala Ala Met Ile Lys Lys Met Cys Pro Ser Asp Ser Glu
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             20                  25                  30

Ser Lys Val Gly Lys Thr Ala Ile Val Ser Arg Phe Leu Thr Gly Arg
         35                  40                  45

Phe Glu Asp Ala Tyr Thr Pro Thr Ile Glu Asp Phe His Arg Lys Phe
     50                  55                  60

Tyr Ser Ile Arg Gly Glu Val Tyr Gln Leu Asp Ile Leu Asp Thr Ser
 65                  70                  75                  80

Gly Asn His Pro Phe Pro Ala Met Arg Arg Leu Ser Ile Leu Thr Gly
                 85                  90                  95

Asp Val Phe Ile Leu Val Phe Ser Leu Asp Asn Arg Asp Ser Phe Glu
            100                 105                 110
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Glu Val Gln Arg Leu Arg Gln Gln Ile Leu Asp Thr Lys Ser Cys Leu
115 120 125

Lys Asn Lys Thr Lys Glu Asn Val Asp Val Pro Leu Val Ile Cys Gly
130 135 140

Asn Lys Gly Asp Arg Asp Phe Tyr Arg Glu Val Asp Gln Arg Glu Ile
145 150 155 160

Glu Gln Leu Val Gly Asp Asp Pro Gln Arg Cys Ala Tyr Phe Glu Ile
165 170 175

Ser Ala Lys Lys Asn Ser Ser Leu Asp Gln Met Phe Arg Ala Leu Phe
180 185 190

Ala Met Ala Lys Leu Pro Ser Glu Met Ser Pro Asp Leu His Arg Lys
195 200 205

Val Ser Val Gln Tyr Cys Asp Val Leu His Lys Lys Ala Leu Arg Asn
210 215 220

Lys Lys Leu Leu Arg Ala Gly Ser Gly Gly Gly Gly Gly Asp Pro Gly
225 230 235 240

Asp Ala Phe Gly Ile Val Ala Pro Phe Ala Arg Arg Pro Ser Val His
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Ser Asp Leu Met Tyr Ile Arg Glu Lys Ala Ser Ala Gly Ser Gln Ala
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Lys Asp Lys Glu Arg Cys Val Ile Ser
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gccccgcgcc cgccctcgcg gcccctctgc cca atg aaa ctg gcc gcg atg atc 174
Met Lys Leu Ala Ala Met Ile
1 5

aag aag atg tgc ccg agc gac tcg gag ctg agt atc ccg gcc aag aac 222
Lys Lys Met Cys Pro Ser Asp Ser Glu Leu Ser Ile Pro Ala Lys Asn
10 15 20

tgc tat cgc atg gtc atc ctc ggc tcg tcc aag gtg ggc aag acg gcc 270
Cys Tyr Arg Met Val Ile Leu Gly Ser Ser Lys Val Gly Lys Thr Ala
25 30 35

atc gtg tcg cgc ttc ctc acc ggc cgc ttc gag gac gcc tac acg cct 318

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Ile Val Ser Arg Phe Leu Thr Gly Arg Phe Glu Asp Ala Tyr Thr Pro
 40                  45                  50                  55

acc atc gag gac ttc cac cgc aag ttc tac tcc atc cgc ggc gag gtc   366
Thr Ile Glu Asp Phe His Arg Lys Phe Tyr Ser Ile Arg Gly Glu Val
                    60                  65                  70

tac cag ctc gac atc ctc gac acg tcc ggc aac cac ccg ttc ccc gcc   414
Tyr Gln Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe Pro Ala
                75                  80                  85

atg cgg cgc ctc tcc atc ctc aca gga gac gtt ttc atc ctg gtg ttc   462
Met Arg Arg Leu Ser Ile Leu Thr Gly Asp Val Phe Ile Leu Val Phe
            90                  95                  100

agt ctg gac aac cgc gac tcc ttc gag gag gtg cag cgg ctc agg cag   510
Ser Leu Asp Asn Arg Asp Ser Phe Glu Glu Val Gln Arg Leu Arg Gln
        105                 110                 115

cag atc ctc gac acc aag tct tgc ctc aag aac aaa acc aag gag aac   558
Gln Ile Leu Asp Thr Lys Ser Cys Leu Lys Asn Lys Thr Lys Glu Asn
120                 125                 130                 135

gtg gac gtg ccc ctg gtc atc tgc ggc aac aag ggt gac cgc gac ttc   606
Val Asp Val Pro Leu Val Ile Cys Gly Asn Lys Gly Asp Arg Asp Phe
                140                 145                 150

tac cgc gag gtg gac cag cgc gag atc gag cag ctg gtg ggc gac gac   654
Tyr Arg Glu Val Asp Gln Arg Glu Ile Glu Gln Leu Val Gly Asp Asp
            155                 160                 165

ccc cag cgc tgc gcc tac ttc gag atc tcg gcc aag aag aac agc agc   702
Pro Gln Arg Cys Ala Tyr Phe Glu Ile Ser Ala Lys Lys Asn Ser Ser
        170                 175                 180

ctg gac cag atg ttc cgc gcg ctc ttc gcc atg gcc aag ctg ccc agc   750
Leu Asp Gln Met Phe Arg Ala Leu Phe Ala Met Ala Lys Leu Pro Ser
    185                 190                 195

gag atg agc cca gac ctg cac cgc aag gtc tcg gtg cag tac tgc gac   798
Glu Met Ser Pro Asp Leu His Arg Lys Val Ser Val Gln Tyr Cys Asp
200                 205                 210                 215

gtg ctg cac aag aag gcg ctg cgg aac aag aag ctg ctg cgg gcc ggc   846
Val Leu His Lys Lys Ala Leu Arg Asn Lys Lys Leu Leu Arg Ala Gly
                220                 225                 230

agc ggc ggc ggc ggc ggc gac ccg ggc gac gcc ttt ggc atc gtg gca   894
Ser Gly Gly Gly Gly Gly Asp Pro Gly Asp Ala Phe Gly Ile Val Ala
            235                 240                 245

ccc ttc gcg cgc cgg ccc agc gta cac agc gac ctc atg tac atc cgc   942
Pro Phe Ala Arg Arg Pro Ser Val His Ser Asp Leu Met Tyr Ile Arg
        250                 255                 260

gag aag gcc agc gcc ggc agc cag gcc aag gac aag gag cgc tgc gtc   990
Glu Lys Ala Ser Ala Gly Ser Gln Ala Lys Asp Lys Glu Arg Cys Val
    265                 270                 275

atc agc taggagcccc gccgcgctgg cgacacaacc taaggaggac ctttttgtta   1046
Ile Ser
280
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agtcaaatcc aacggcccgg tgcgccccag gccgggagcg cgcgcggact ggcgtctccc 1106

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attgaggccc gccacccccа taactttggg agcgagggcc cagccgaggg tggatttatc 1286

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gctaggcttc aagaaaccgt catgcccgct tgagggtcag gacccacggg gcattatctt 1406

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gggggggcgtg ggtcaaatca tagccaagtg acttgtttac atgtgagtga aactgcacaa 1526

aggaacacaa aacaaaactt gcactttaac ggtagttccg gtgtcaacat ggacacgaac 1586

aaaaccttac ccaggtgttt atactgtgtg tgtgtgaggt ctttaaagtt attgctttat 1646

ttggtttttt aatatacaat aaaataattt aaaatggaaa aaaaaaaaaa aaaaaaaaaa 1706

aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aagcggccgc tcgagcatgc 1766

atctagaggg ccgcatcatg taattagtta tgaac 1801

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<212> DNA
<213> Artificial sequence
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<400> 6
gtttgatgtg ggtgctcagc ggtctgag 28

<210> 7
<211> 29

<212> DNA
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ctcagaaccg ctgagcaccc acatcaaac 29

<210> 8
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cgggatccat gaaactggcc gcgatgatca agaag 35

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ggatccatgc aaacgctaaa gtgtg 25

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<210> 12
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<400> 13
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<213> Artificial sequence
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<400> 16
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<210> 17
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<213> Artificial sequence
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<400> 17
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<210> 18
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<212> PRT
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<220> Xaa's at positions 2,3 may be any amino acid

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<210> 20
<211> 4
<212> PRT
<213> Unknown

<220> Xaa at position 3 may be any amino acid

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Asn Lys Xaa Asp
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<210> 21
<211> 5
<212> PRT
<213> Unknown

<220> Xaa at position 2 may be any amino acid

<400> 21
Glu Xaa Ser Ala Lys
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<210> 22
<211> 4
<212> PRT
<213> Unknown

<220> Xaa at position 2 or 3 may be any aliphatic amino acid residue
<220> Xaa at position 4 may be any amino acid

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<222> (45)..(587)

<400> 24

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ctt gca gct ttg ccc cac tca tgc ctg gaa agg gcc aag gag att aag   104
Leu Ala Ala Leu Pro His Ser Cys Leu Glu Arg Ala Lys Glu Ile Lys
  5                  10                  15                  20

atc aag ttg gga att ctc ctc cag aag cca gac tca gtt ggt gac ctt   152
Ile Lys Leu Gly Ile Leu Leu Gln Lys Pro Asp Ser Val Gly Asp Leu
                  25                  30                  35

gtc att ccg tac aat gag aag cca gag aaa cca gcc aag acc cag aaa   200
Val Ile Pro Tyr Asn Glu Lys Pro Glu Lys Pro Ala Lys Thr Gln Lys
              40                  45                  50

acc tcg ctg gac gag gcc ctg cag tgg cgt gat tcc ctg gac aaa ctc   248
Thr Ser Leu Asp Glu Ala Leu Gln Trp Arg Asp Ser Leu Asp Lys Leu
          55                  60                  65

ctg cag aac aac tat gga ctt gcc agt ttc aaa agt ttc ctg aag tct   296
Leu Gln Asn Asn Tyr Gly Leu Ala Ser Phe Lys Ser Phe Leu Lys Ser
      70                  75                  80

gaa ttc agt gag gaa aac ctt gag ttc tgg att gcc tgt gag gat tac   344
Glu Phe Ser Glu Glu Asn Leu Glu Phe Trp Ile Ala Cys Glu Asp Tyr
  85                  90                  95                 100

aag aag atc aag tcc cct gcc aag atg gct gag aag gca aag caa att   392
Lys Lys Ile Lys Ser Pro Ala Lys Met Ala Glu Lys Ala Lys Gln Ile
                 105                 110                 115

tat gaa gaa ttc att caa acg gag gct cct aaa gag gtg aat att gac   440
Tyr Glu Glu Phe Ile Gln Thr Glu Ala Pro Lys Glu Val Asn Ile Asp
             120                 125                 130

cac ttc act aag gac atc aca atg aag aac ctg gtg gaa cct tcc ctg   488
His Phe Thr Lys Asp Ile Thr Met Lys Asn Leu Val Glu Pro Ser Leu
         135                 140                 145

agc agc ttt gac atg gcc cag aaa aga atc cat gcc ctg atg gaa aag   536
Ser Ser Phe Asp Met Ala Gln Lys Arg Ile His Ala Leu Met Glu Lys
     150                 155                 160

gat tct ctg cct cgc ttt gtg cgc tct gag ttt tat cag gag tta atc   584
Asp Ser Leu Pro Arg Phe Val Arg Ser Glu Phe Tyr Gln Glu Leu Ile
165                 170                 175                 180

aag tagtaattta gccaggctat gaaatcatcc tgtgagttat ttcctccata       637
Lys

ataaccctgc atttcccatt aatctacata tcttcccaca gcagctttgc tcagtgatac 697

ccacatggga aaaatcccag gggatgttgc ttactctttt tgcccacact gctttggata 757

cttatctact gtccgaaggc cttctttccc cactcaattc ttcctgccct gttattaatt 817
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ctaaattggt ctaaaaaaga atattaagtg tggacagacc tatttcaaag gagcttaatt 997

gatctcactt gttttagttc tgatccaggg agatcacccc tctaattatt tctgaacttg 1057

gttaataaaa gtttataaga tttttatgaa gcagccactg tatgatattt taagcaaata 1117

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<213> Homo sapiens

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Met Cys Lys Gly Leu Ala Ala Leu Pro His Ser Cys Leu Glu Arg Ala
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Lys Glu Ile Lys Ile Lys Leu Gly Ile Leu Leu Gln Lys Pro Asp Ser
            20                  25                  30

Val Gly Asp Leu Val Ile Pro Tyr Asn Glu Lys Pro Glu Lys Pro Ala
        35                  40                  45

Lys Thr Gln Lys Thr Ser Leu Asp Glu Ala Leu Gln Trp Arg Asp Ser
    50                  55                  60

Leu Asp Lys Leu Leu Gln Asn Asn Tyr Gly Leu Ala Ser Phe Lys Ser
65                  70                  75                  80

Phe Leu Lys Ser Glu Phe Ser Glu Glu Asn Leu Glu Phe Trp Ile Ala
                85                  90                  95

Cys Glu Asp Tyr Lys Lys Ile Lys Ser Pro Ala Lys Met Ala Glu Lys
            100                 105                 110

Ala Lys Gln Ile Tyr Glu Glu Phe Ile Gln Thr Glu Ala Pro Lys Glu
        115                 120                 125

Val Asn Ile Asp His Phe Thr Lys Asp Ile Thr Met Lys Asn Leu Val
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130 135 140

Glu Pro Ser Leu Ser Ser Phe Asp Met Ala Gln Lys Arg Ile His Ala
145 150 155 160

Leu Met Glu Lys Asp Ser Leu Pro Arg Phe Val Arg Ser Glu Phe Tyr
165 170 175

Gln Glu Leu Ile Lys
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<210> 28
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<210> 29
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<210> 31

<211> 34
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<210> 33
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  1               5                  10                  15

ttg tcc agc ggg aac tgc acg ctc agt gtg ccc gcc aaa aac tca tac   96
Leu Ser Ser Gly Asn Cys Thr Leu Ser Val Pro Ala Lys Asn Ser Tyr
             20                  25                  30

cgc atg gtg gtg ctg ggt gcc tct cgg gtg ggc aag agc tcc atc gtg   144
Arg Met Val Val Leu Gly Ala Ser Arg Val Gly Lys Ser Ser Ile Val
         35                  40                  45

tct cgc ttc ctc aat ggc cgc ttt gag gac cag tac aca ccc acc atc   192
Ser Arg Phe Leu Asn Gly Arg Phe Glu Asp Gln Tyr Thr Pro Thr Ile
     50                  55                  60

gag gac ttc cac cgt aag gta tac aac atc cgc ggc gac atg tac cag   240
Glu Asp Phe His Arg Lys Val Tyr Asn Ile Arg Gly Asp Met Tyr Gln
 65                  70                  75                  80

ctc gac atc ctg gat acc tct ggc aac cac ccc ttc ccc gcc atg cgc   288
Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe Pro Ala Met Arg
                 85                  90                  95

agg ctg tcc atc ctc aca ggg gat gtc ttc atc ctg gtg ttc agc ctg   336
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Arg Leu Ser Ile Leu Thr Gly Asp Val Phe Ile Leu Val Phe Ser Leu
            100                 105                 110

gat aac cgg gag tcc ttc gat gag gtc aag cgc ctt cag aag cag atc   384
Asp Asn Arg Glu Ser Phe Asp Glu Val Lys Arg Leu Gln Lys Gln Ile
        115                 120                 125

ctg gag gtc aag tcc tgc ctg aag aac aag acc aag gag gcg gcg gag   432
Leu Glu Val Lys Ser Cys Leu Lys Asn Lys Thr Lys Glu Ala Ala Glu
    130                 135                 140

ctg ccc atg gtc atc tgt ggc aac aag aac gac cac ggc gag ctg tgc   480
Leu Pro Met Val Ile Cys Gly Asn Lys Asn Asp His Gly Glu Leu Cys
145                 150                 155                 160

cgc cag gtg ccc acc acc gag gcc gag ctg ctg gtg tcg ggc gac gag   528
Arg Gln Val Pro Thr Thr Glu Ala Glu Leu Leu Val Ser Gly Asp Glu
                165                 170                 175

aac tgc gcc tac ttc gag gtg tcg gcc aag aag aac acc aac gtg gac   576
Asn Cys Ala Tyr Phe Glu Val Ser Ala Lys Lys Asn Thr Asn Val Asp
            180                 185                 190

gag atg ttc tac gtg ctc ttc agc atg gcc aag ctg cca cac gag atg   624
Glu Met Phe Tyr Val Leu Phe Ser Met Ala Lys Leu Pro His Glu Met
        195                 200                 205

agc ccc gcc ctg cat cgc aag atc tcc gtg cag tac ggt gac gcc ttc   672
Ser Pro Ala Leu His Arg Lys Ile Ser Val Gln Tyr Gly Asp Ala Phe
    210                 215                 220

cac ccc agg ccc ttc tgc atg cgc cgc gtc aag gag atg gac gcc tat   720
His Pro Arg Pro Phe Cys Met Arg Arg Val Lys Glu Met Asp Ala Tyr
225                 230                 235                 240

ggc atg gtc tcg ccc ttc gcc cgc cgc ccc agc gtc aac agt gac ctc   768
Gly Met Val Ser Pro Phe Ala Arg Arg Pro Ser Val Asn Ser Asp Leu
                245                 250                 255

aag tac atc aag gcc aag gtc ctt cgg gaa ggc cag gcc cgt gag agg   816
Lys Tyr Ile Lys Ala Lys Val Leu Arg Glu Gly Gln Ala Arg Glu Arg
            260                 265                 270

gac aag tgc acc atc cag tga                                      837
Asp Lys Cys Thr Ile Gln
        275


<210> 41
<211> 278
<212> PRT
<213> Homo sapiens

<400> 41
Met Pro Ala Ser Leu Ala Leu Leu Gln Pro Arg Ala Met Met Lys Thr
  1               5                  10                  15

Leu Ser Ser Gly Asn Cys Thr Leu Ser Val Pro Ala Lys Asn Ser Tyr
            20                  25                  30

Arg Met Val Val Leu Gly Ala Ser Arg Val Gly Lys Ser Ser Ile Val
        35                  40                  45
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Ser Arg Phe Leu Asn Gly Arg Phe Glu Asp Gln Tyr Thr Pro Thr Ile
50 55 60

Glu Asp Phe His Arg Lys Val Tyr Asn Ile Arg Gly Asp Met Tyr Gln
65 70 75 80

Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe Pro Ala Met Arg
85 90 95

Arg Leu Ser Ile Leu Thr Gly Asp Val Phe Ile Leu Val Phe Ser Leu
100 105 110

Asp Asn Arg Glu Ser Phe Asp Glu Val Lys Arg Leu Gln Lys Gln Ile
115 120 125

Leu Glu Val Lys Ser Cys Leu Lys Asn Lys Thr Lys Glu Ala Ala Glu
130 135 140

Leu Pro Met Val Ile Cys Gly Asn Lys Asn Asp His Gly Glu Leu Cys
145 150 155 160

Arg Gln Val Pro Thr Thr Glu Ala Glu Leu Leu Val Ser Gly Asp Glu
165 170 175

Asn Cys Ala Tyr Phe Glu Val Ser Ala Lys Lys Asn Thr Asn Val Asp
180 185 190

Glu Met Phe Tyr Val Leu Phe Ser Met Ala Lys Leu Pro His Glu Met
195 200 205

Ser Pro Ala Leu His Arg Lys Ile Ser Val Gln Tyr Gly Asp Ala Phe
210 215 220

His Pro Arg Pro Phe Cys Met Arg Arg Val Lys Glu Met Asp Ala Tyr
225 230 235 240

Gly Met Val Ser Pro Phe Ala Arg Arg Pro Ser Val Asn Ser Asp Leu
245 250 255

Lys Tyr Ile Lys Ala Lys Val Leu Arg Glu Gly Gln Ala Arg Glu Arg
260 265 270

Asp Lys Cys Thr Ile Gln
275

<210> 42
<211> 15
<212> PRT
<213> Homo sapiens

<400> 42
Asp Thr Lys Ser Cys Leu Lys Asn Lys Thr Lys Glu Asn Val Asp
1 5 10 15

<210> 43
<211> 19
<212> DNA
<213> Artificial sequence
<223> Probe/Primer

<400> 43
ttctcgcgga tgtacatga 19

<210> 44
<211> 20
<212> DNA
<213> Artificial sequence
<223> Probe/Primer

<400> 44
tccaccgcaa gttctactcc 20

<210> 45
<211> 1740
<212> DNA
<213> Homo sapiens

<220>
<221> CDS
<222> (146)..(988)

<400> 45
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tcaccccgcg tgccacccca gcgaccctca gccgctctct gcccttctct cggccccgcg 120

cccgccctcg cggcccctct gccca atg aaa ctg gcc gcg atg atc aag aag 172
Met Lys Leu Ala Ala Met Ile Lys Lys
1 5

atg tgc ccg agc gac tcg gag ctg agt atc ccg gcc aag aac tgc tat 220
Met Cys Pro Ser Asp Ser Glu Leu Ser Ile Pro Ala Lys Asn Cys Tyr
10 15 20 25

cgc atg gtc atc ctc ggc tcg tcc aag gtg ggc aag acg gcc atc gtg 268
Arg Met Val Ile Leu Gly Ser Ser Lys Val Gly Lys Thr Ala Ile Val
30 35 40

tcg cgc ttc ctc acc ggc cgc ttc gag gac gcc tac acg cct acc atc 316
Ser Arg Phe Leu Thr Gly Arg Phe Glu Asp Ala Tyr Thr Pro Thr Ile
45 50 55

gag gac ttc cac cgc aag ttc tac tcc atc cgc ggc gag gtc tac cag 364
Glu Asp Phe His Arg Lys Phe Tyr Ser Ile Arg Gly Glu Val Tyr Gln
60 65 70

ctc gac atc ctc gac acg tcc ggc aac cac ccg ttc ccc gcc atg cgg 412
Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe Pro Ala Met Arg
75 80 85

cgc ctc tcc atc ctc aca gga gac gtt ttc atc ctg gtg ttc agt ctg 460
Arg Leu Ser Ile Leu Thr Gly Asp Val Phe Ile Leu Val Phe Ser Leu
90 95 100 105

gac aac cgc gac tcc ttc gag gag gtg cag cgg ctc agg cag cag atc 508
Asp Asn Arg Asp Ser Phe Glu Glu Val Gln Arg Leu Arg Gln Gln Ile
110 115 120

ctc gac acc aag tct tgc ctc aag aac aaa acc aag gag aac gtg gac 556

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Leu Asp Thr Lys Ser Cys Leu Lys Asn Lys Thr Lys Glu Asn Val Asp
            125                 130                 135

gtg ccc ctg gtc atc tgc ggc aac aag ggt gac cgc gac ttc tac cgc   604
Val Pro Leu Val Ile Cys Gly Asn Lys Gly Asp Arg Asp Phe Tyr Arg
        140                 145                 150

gag gtg gac cag cgc gag atc gag cag ctg gtg ggc gac gac ccc cag   652
Glu Val Asp Gln Arg Glu Ile Glu Gln Leu Val Gly Asp Asp Pro Gln
    155                 160                 165

cgc tgc gcc tac ttc gag atc tcg gcc aag aag aac agc agc ctg gac   700
Arg Cys Ala Tyr Phe Glu Ile Ser Ala Lys Lys Asn Ser Ser Leu Asp
170                 175                 180                 185

cag atg ttc cgc gcg ctc ttc gcc atg gcc aag ctg ccc agc gag atg   748
Gln Met Phe Arg Ala Leu Phe Ala Met Ala Lys Leu Pro Ser Glu Met
                190                 195                 200

agc cca gac ctg cac cgc aag gtc tcg gtg cag tac tgc gac gtg ctg   796
Ser Pro Asp Leu His Arg Lys Val Ser Val Gln Tyr Cys Asp Val Leu
            205                 210                 215

cac aag aag gcg ctg cgg aac aag aag ctg ctg cgg gcc ggc agc ggc   844
His Lys Lys Ala Leu Arg Asn Lys Lys Leu Leu Arg Ala Gly Ser Gly
        220                 225                 230

ggc ggc ggc ggc gac ccg ggc gac gcc ttt ggc atc gtg gca ccc ttc   892
Gly Gly Gly Gly Asp Pro Gly Asp Ala Phe Gly Ile Val Ala Pro Phe
    235                 240                 245

gcg cgc cgg ccc agc gta cac agc gac ctc atg tac atc cgc gag aag   940
Ala Arg Arg Pro Ser Val His Ser Asp Leu Met Tyr Ile Arg Glu Lys
250                 255                 260                 265

gcc agc gcc ggc agc cag gcc aag gac aag gag cgc tgc gtc atc agc   988
Ala Ser Ala Gly Ser Gln Ala Lys Asp Lys Glu Arg Cys Val Ile Ser
                270                 275                 280

taggagcccc gccgcgctgg cgacacaacc taaggaggac ctttttgtta agtcaaatcc 1048

aacggcccgg tgcgccccag gccgggagcg cgcgcggact ggcgtctccc ctcccggcga 1108

tccgccccca gcactgggga ggcgccactg aaccgagaag ggacggtcat ctgctccgga 1168

aggaaagaga acgggccaag actgggacta ttccccaccc ccggtccccc attgaggccc 1228

gccaccccca taactttggg agcgagggcc cagccgaggg tggatttatc ttctcaaaga 1288

cctaagagtg agcgcggggt gggggaggga tgtgaagtta tccagcctct gctaggcttc 1348

aagaaaccgt catgcccgct tgagggtcag gacccacggg gcattatctt gtctgtgatt 1408

ccgggttgct gtgacagccg gtagagcctc tgccctcccg aaactaagcg gggggggcgtg 1468

ggtcaaatca tagccaagtg acttgtttac atgtgagtga aactgcacaa aggaacacaa 1528

aacaaaactt gcactttaac ggtagttccg gtgtcaacat ggacacgaac aaaaccttac 1588

ccaggtgttt atactgtgtg tgtgtgaggt ctttaaagtt attgctttat ttggtttttt 1648

aatatacaat aaaataattt aaaatggaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 1708
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aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aa 1740

<210> 46
<211> 281
<212> PRT
<213> Homo sapiens

<400> 46

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Met Lys Leu Ala Ala Met Ile Lys Lys Met Cys Pro Ser Asp Ser Glu
  1               5                  10                  15

Leu Ser Ile Pro Ala Lys Asn Cys Tyr Arg Met Val Ile Leu Gly Ser
             20                  25                  30

Ser Lys Val Gly Lys Thr Ala Ile Val Ser Arg Phe Leu Thr Gly Arg
         35                  40                  45

Phe Glu Asp Ala Tyr Thr Pro Thr Ile Glu Asp Phe His Arg Lys Phe
     50                  55                  60

Tyr Ser Ile Arg Gly Glu Val Tyr Gln Leu Asp Ile Leu Asp Thr Ser
 65                  70                  75                  80

Gly Asn His Pro Phe Pro Ala Met Arg Arg Leu Ser Ile Leu Thr Gly
                 85                  90                  95

Asp Val Phe Ile Leu Val Phe Ser Leu Asp Asn Arg Asp Ser Phe Glu
            100                 105                 110

Glu Val Gln Arg Leu Arg Gln Gln Ile Leu Asp Thr Lys Ser Cys Leu
        115                 120                 125

Lys Asn Lys Thr Lys Glu Asn Val Asp Val Pro Leu Val Ile Cys Gly
    130                 135                 140

Asn Lys Gly Asp Arg Asp Phe Tyr Arg Glu Val Asp Gln Arg Glu Ile
145                 150                 155                 160

Glu Gln Leu Val Gly Asp Asp Pro Gln Arg Cys Ala Tyr Phe Glu Ile
                165                 170                 175

Ser Ala Lys Lys Asn Ser Ser Leu Asp Gln Met Phe Arg Ala Leu Phe
            180                 185                 190

Ala Met Ala Lys Leu Pro Ser Glu Met Ser Pro Asp Leu His Arg Lys
        195                 200                 205

Val Ser Val Gln Tyr Cys Asp Val Leu His Lys Lys Ala Leu Arg Asn
    210                 215                 220

Lys Lys Leu Leu Arg Ala Gly Ser Gly Gly Gly Gly Gly Asp Pro Gly
225                 230                 235                 240

Asp Ala Phe Gly Ile Val Ala Pro Phe Ala Arg Arg Pro Ser Val His
                245                 250                 255

Ser Asp Leu Met Tyr Ile Arg Glu Lys Ala Ser Ala Gly Ser Gln Ala
            260                 265                 270

Lys Asp Lys Glu Arg Cys Val Ile Ser
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275 280

<210> 47
<211> 189
<212> PRT
<213> Homo sapiens

<400> 47
Met Thr Glu Tyr Lys Leu Val Val Val Gly Ala Gly Gly Val Gly Lys
1 5 10 15

Ser Ala Leu Thr Ile Gln Leu Ile Gln Asn His Phe Val Asp Glu Tyr
20 25 30

Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gln Val Val Ile Asp Gly
35 40 45

Glu Thr Cys Leu Leu Asp Ile Leu Asp Thr Ala Gly Gln Glu Glu Tyr
50 55 60

Ser Ala Met Arg Asp Gln Tyr Met Arg Thr Gly Glu Gly Phe Leu Cys
65 70 75 80

Val Phe Ala Ile Asn Asn Thr Lys Ser Phe Glu Asp Ile His Gln Tyr
85 90 95

Arg Glu Gln Ile Lys Arg Val Lys Asp Ser Asp Asp Val Pro Met Val
100 105 110

Leu Val Gly Asn Lys Cys Asp Leu Ala Ala Arg Thr Val Glu Ser Arg
115 120 125

Gln Ala Gln Asp Leu Ala Arg Ser Tyr Gly Ile Pro Tyr Ile Glu Thr
130 135 140

Ser Ala Lys Thr Arg Gln Gly Val Glu Asp Ala Phe Tyr Thr Leu Val
145 150 155 160

Arg Glu Ile Arg Gln His Lys Leu Arg Lys Leu Asn Pro Pro Asp Glu
165 170 175

Ser Gly Pro Gly Cys Met Ser Cys Lys Cys Val Leu Ser
180 185

<210> 48
<211> 206
<212> PRT
<213> Homo sapiens

<400> 48
Met Ala Ala Asn Lys Pro Lys Gly Gln Asn Ser Leu Ala Leu His Lys
1 5 10 15

Val Ile Met Val Gly Ser Gly Gly Val Gly Lys Ser Ala Leu Thr Leu
20 25 30

Gln Phe Met Tyr Asp Glu Phe Val Glu Asp Tyr Glu Pro Thr Lys Ala
35 40 45

Asp Ser Tyr Arg Lys Lys Val Val Leu Asp Gly Glu Glu Val Gln Ile
50 55 60

Asp Ile Leu Asp Thr Ala Gly Gln Glu Asp Tyr Ala Ala Ile Arg Asp
65 70 75 80

Asn Tyr Phe Arg Ser Gly Glu Gly Phe Leu Cys Val Phe Ser Ile Thr
85 90 95

Glu Met Glu Ser Phe Ala Ala Thr Ala Asp Phe Arg Glu Gln Ile Leu
100 105 110

Arg Val Lys Glu Asp Glu Asn Val Pro Phe Leu Leu Val Gly Asn Lys
115 120 125

Ser Asp Leu Glu Asp Lys Arg Gln Val Ser Val Glu Glu Ala Lys Asn
130 135 140

Arg Ala Glu Gln Trp Asn Val Asn Tyr Val Glu Thr Ser Ala Lys Thr
145 150 155 160

Arg Ala Asn Val Asp Lys Val Phe Phe Asp Leu Met Arg Glu Ile Arg
165 170 175

Ala Arg Lys Met Glu Asp Ser Lys Glu Lys Asn Gly Lys Lys Lys Arg
180 185 190

Lys Ser Leu Ala Lys Arg Ile Arg Glu Arg Cys Cys Ile Leu
195 200 205

<210> 49
<211> 205
<212> PRT
<213> Homo sapiens

<400> 49
Met Ser Ser Met Asn Pro Glu Tyr Asp Tyr Leu Phe Lys Leu Leu Leu
1 5 10 15

Ile Gly Asp Ser Gly Val Gly Lys Ser Cys Leu Leu Leu Arg Phe Ala
20 25 30

Asp Asp Thr Tyr Thr Glu Ser Tyr Ile Ser Thr Ile Gly Val Asp Phe
35 40 45

Lys Ile Arg Thr Ile Glu Leu Asp Gly Lys Thr Ile Lys Leu Gln Ile
50 55 60

Trp Asp Thr Ala Gly Gln Glu Arg Phe Arg Thr Ile Thr Ser Ser Tyr
65 70 75 80

Tyr Arg Gly Ala His Gly Ile Ile Val Val Tyr Asp Val Thr Asp Gln
85 90 95

Glu Ser Phe Asn Asn Val Lys Gln Trp Leu Gln Glu Ile Asp Arg Tyr
100 105 110

Ala Ser Glu Asn Val Asn Lys Leu Leu Val Gly Asn Lys Cys Asp Leu
115 120 125

Thr Thr Lys Lys Val Val Asp Tyr Thr Thr Ala Lys Glu Phe Ala Asp
130 135 140

Ser Leu Gly Ile Pro Phe Leu Glu Thr Ser Ala Lys Asn Ala Thr Asn
145 150 155 160

Val Glu Gln Ser Phe Met Thr Met Ala Ala Glu Ile Lys Lys Arg Met
165 170 175

Gly Pro Gly Ala Thr Ala Gly Gly Ala Glu Lys Ser Asn Val Lys Ile
180 185 190

Gln Ser Thr Pro Val Lys Gln Ala Gly Gly Gly Cys Cys
195 200 205

<210> 50
<211> 210
<212> PRT
<213> Homo sapiens

<400> 50
Met Thr Ala Ala Gln Ala Ala Gly Glu Glu Ala Pro Pro Gly Val Arg
1 5 10 15

Ser Val Lys Val Val Leu Val Gly Asp Gly Gly Cys Gly Lys Thr Ser
20 25 30

Leu Leu Met Val Phe Ala Asp Gly Ala Phe Pro Glu Ser Tyr Thr Pro
35 40 45

Thr Val Phe Glu Arg Tyr Met Val Asn Leu Gln Val Lys Gly Lys Pro
50 55 60

Val His Leu His Ile Trp Asp Thr Ala Gly Gln Asp Asp Tyr Asp Arg
65 70 75 80

Leu Arg Pro Leu Phe Tyr Pro Asp Ala Ser Val Leu Leu Leu Cys Phe
85 90 95

Asp Val Thr Ser Pro Asn Ser Phe Asp Asn Ile Phe Asn Arg Trp Tyr
100 105 110

Pro Glu Val Asn His Phe Cys Lys Lys Val Pro Ile Ile Val Val Gly
115 120 125

Cys Lys Thr Asp Leu Arg Lys Asp Lys Ser Leu Val Asn Lys Leu Arg
130 135 140

Arg Asn Gly Leu Glu Pro Val Thr Tyr His Arg Gly Gln Glu Met Ala
145 150 155 160

Arg Ser Val Gly Ala Val Ala Tyr Leu Glu Cys Ser Ala Arg Leu His
165 170 175

Asp Asn Val His Ala Val Phe Gln Glu Ala Ala Glu Val Ala Leu Ser
180 185 190

Ser Arg Gly Arg Asn Phe Trp Arg Arg Ile Thr Gln Gly Phe Cys Val
195 200 205

Val Thr
210

<210> 51
<211> 191
<212> PRT
<213> Homo sapiens

<400> 51
Met Gln Thr Ile Lys Cys Val Val Val Gly Asp Gly Ala Val Gly Lys
1 5 10 15

Thr Cys Leu Leu Ile Ser Tyr Thr Thr Asn Lys Phe Pro Ser Glu Tyr
20 25 30

Val Pro Thr Val Phe Asp Asn Tyr Ala Val Thr Val Met Ile Gly Gly
35 40 45

Glu Pro Tyr Thr Leu Gly Leu Phe Asp Thr Ala Gly Gln Glu Asp Tyr
50 55 60

Asp Arg Leu Arg Pro Leu Ser Tyr Pro Gln Thr Asp Val Phe Leu Val
65 70 75 80

Cys Phe Ser Val Val Ser Pro Ser Ser Phe Glu Asn Val Lys Glu Lys
85 90 95

Trp Val Pro Glu Ile Thr His His Cys Pro Lys Thr Pro Phe Leu Leu
100 105 110

Val Gly Thr Gln Ile Asp Leu Arg Asp Asp Pro Ser Thr Ile Glu Lys
115 120 125

Leu Ala Lys Asn Lys Gln Lys Pro Ile Thr Pro Glu Thr Ala Glu Lys
130 135 140

Leu Ala Arg Asp Leu Lys Ala Val Lys Tyr Val Glu Cys Ser Ala Leu
145 150 155 160

Thr Gln Arg Gly Leu Lys Asn Val Phe Asp Glu Ala Ile Leu Ala Ala
165 170 175

Leu Glu Pro Pro Glu Thr Gln Pro Lys Arg Lys Cys Cys Ile Phe
180 185 190

<210> 52
<211> 192
<212> PRT
<213> Homo sapiens

<400> 52
Met Gln Ala Ile Lys Cys Val Val Val Gly Asp Gly Ala Val Gly Lys
1 5 10 15

Thr Cys Leu Leu Ile Ser Tyr Thr Thr Asn Ala Phe Pro Gly Glu Tyr
20 25 30

Ile Pro Thr Val Phe Asp Asn Tyr Ser Ala Asn Val Met Val Asp Ser
35 40 45

Lys Pro Val Asn Leu Gly Leu Trp Asp Thr Ala Gly Gln Glu Asp Tyr
50 55 60

Asp Arg Leu Arg Pro Leu Ser Tyr Pro Gln Thr Asp Val Phe Leu Ile

65 70 75 80

Cys Phe Ser Leu Val Ser Pro Ala Ser Tyr Glu Asn Val Arg Ala Lys
85 90 95

Trp Phe Pro Glu Val Arg His His Cys Pro Ser Thr Pro Ile Ile Leu
100 105 110

Val Gly Thr Lys Leu Asp Leu Arg Asp Asp Lys Asp Thr Ile Glu Lys
115 120 125

Leu Lys Glu Lys Lys Leu Ala Pro Ile Thr Tyr Pro Gln Gly Leu Ala
130 135 140

Leu Ala Lys Glu Ile Asp Ser Val Lys Tyr Leu Glu Cys Ser Ala Leu
145 150 155 160

Thr Gln Arg Gly Leu Lys Thr Val Phe Asp Glu Ala Ile Arg Ala Val
165 170 175

Leu Cys Pro Gln Pro Thr Arg Gln Gln Lys Arg Ala Cys Ser Leu Leu
180 185 190

<210> 53
<211> 181
<212> PRT
<213> Homo sapiens

<400> 53
Met Gly Gly Phe Phe Ser Ser Ile Phe Ser Ser Leu Phe Gly Thr Arg
1 5 10 15

Glu Met Arg Ile Leu Ile Leu Gly Leu Asp Gly Ala Gly Lys Thr Thr
20 25 30

Ile Leu Tyr Arg Leu Gln Val Gly Glu Val Val Thr Thr Ile Pro Thr
35 40 45

Ile Gly Phe Asn Val Glu Thr Val Thr Tyr Lys Asn Leu Lys Phe Gln
50 55 60

Val Trp Asp Leu Gly Gly Gln Thr Ser Ile Arg Pro Tyr Trp Arg Cys
65 70 75 80

Tyr Tyr Ser Asn Thr Asp Ala Val Ile Tyr Val Val Asp Ser Cys Asp
85 90 95

Arg Asp Arg Ile Gly Ile Ser Lys Ser Glu Leu Val Ala Met Leu Glu
100 105 110

Glu Glu Glu Leu Arg Lys Ala Ile Leu Val Val Phe Ala Asn Lys Gln
115 120 125

Asp Met Glu Gln Ala Met Thr Ser Ser Glu Met Ala Asn Ser Leu Gly
130 135 140

Leu Pro Ala Leu Lys Asp Arg Lys Trp Gln Ile Phe Lys Thr Ser Ala
145 150 155 160

Thr Lys Gly Thr Gly Leu Asp Glu Ala Met Glu Trp Leu Val Glu Thr
165 170 175

Leu Lys Ser Arg Gln
180

<210> 54
<211> 229
<212> PRT
<213> Homo sapiens

<400> 54
Met Asp Pro Asn Gln Asn Val Lys Cys Lys Ile Val Val Val Gly Asp
1 5 10 15

Ser Gln Cys Gly Lys Thr Ala Leu Leu His Val Phe Ala Lys Asp Cys
20 25 30

Phe Pro Glu Asn Tyr Val Pro Thr Val Phe Glu Asn Tyr Thr Ala Ser
35 40 45

Phe Glu Ile Asp Thr Gln Arg Ile Glu Leu Ser Leu Trp Asp Thr Ser
50 55 60

Gly Ser Pro Tyr Tyr Asp Asn Val Arg Pro Leu Ser Tyr Pro Asp Ser
65 70 75 80

Asp Ala Val Leu Ile Cys Phe Asp Ile Ser Arg Pro Glu Thr Leu Asp
85 90 95

Ser Val Leu Lys Lys Trp Lys Gly Glu Ile Gln Glu Phe Cys Pro Asn
100 105 110

Thr Lys Met Leu Leu Val Gly Cys Lys Ser Asp Leu Arg Thr Asp Val
115 120 125

Ser Thr Leu Val Glu Leu Ser Asn His Arg Gln Thr Pro Val Ser Tyr
130 135 140

Asp Gln Gly Ala Asn Met Ala Lys Gln Ile Gly Ala Ala Thr Tyr Ile
145 150 155 160

Glu Cys Ser Ala Leu Gln Ser Glu Asn Ser Val Arg Asp Ile Phe His
165 170 175

Val Ala Thr Leu Ala Cys Val Asn Lys Thr Asn Lys Asn Val Lys Arg
180 185 190

Asn Lys Ser Gln Arg Ala Thr Lys Arg Ile Ser His Met Pro Ser Arg
195 200 205

Pro Glu Leu Ser Ala Val Ala Thr Asp Leu Arg Lys Asp Lys Ala Lys
210 215 220

Ser Cys Thr Val Met
225

<210> 55
<211> 16
<212> PRT

<213> Homo sapiens

<400> 55
Lys Ile Val Val Val Gly Asp Ser Gln Cys Gly Lys Thr Ala Leu Leu
1 5 10 15

<210> 56
<211> 16
<212> PRT
<213> Homo sapiens

<400> 56
Lys Ile Val Val Val Gly Asp Ala Glu Cys Gly Lys Thr Ala Leu Leu
1 5 10 15

<210> 57
<211> 16
<212> PRT
<213> Homo sapiens

<400> 57
Lys Leu Val Leu Val Gly Asp Val Gln Cys Gly Lys Thr Ala Met Leu
1 5 10 15

<210> 58
<211> 16
<212> PRT
<213> Homo sapiens

<400> 58
Lys Leu Val Ile Val Gly Asp Gly Ala Cys Gly Lys Thr Cys Leu Leu
1 5 10 15

<210> 59
<211> 16
<212> PRT
<213> Homo sapiens

<400> 59
Lys Leu Val Val Val Gly Asp Gly Ala Cys Gly Lys Thr Cys Leu Leu
1 5 10 15

<210> 60
<211> 16
<212> PRT
<213> Homo sapiens

<400> 60
Lys Cys Val Val Val Gly Asp Gly Ala Val Gly Lys Thr Cys Leu Leu
1 5 10 15

<210> 61
<211> 16
<212> PRT
<213> Homo sapiens

<400> 61
Lys Cys Val Val Val Gly Asp Gly Ala Val Gly Lys Thr Cys Leu Leu
1 5 10 15

<210> 62
<211> 16
<212> PRT
<213> Homo sapiens

<400> 62
Lys Leu Val Val Val Gly Ala Gly Gly Val Gly Lys Ser Ala Leu Thr
1 5 10 15

<210> 63
<211> 16
<212> PRT
<213> Homo sapiens

<400> 63
Arg Met Val Ile Leu Gly Ser Ser Lys Val Gly Lys Thr Ala Ile Val
1 5 10 15

<210> 64
<211> 13
<212> PRT
<213> Homo sapiens

<400> 64
Leu Ser Leu Trp Asp Thr Ser Gly Ser Pro Tyr Tyr Asp
1 5 10

<210> 65
<211> 13
<212> PRT
<213> Homo sapiens

<400> 65
Leu Asn Met Trp Asp Thr Ser Gly Ser Ser Tyr Tyr Asp
1 5 10

<210> 66
<211> 13
<212> PRT
<213> Homo sapiens

<400> 66
Leu Ser Leu Trp Asp Thr Ser Gly Ser Pro Tyr Tyr Asp
1 5 10

<210> 67
<211> 13
<212> PRT
<213> Homo sapiens

<400> 67
Leu Ala Leu Trp Asp Thr Ala Gly Gln Glu Asp Tyr Asp
1 5 10

<210> 68
<211> 13
<212> PRT
<213> Homo sapiens

<400> 68
Leu Ala Leu Trp Asp Thr Ala Gly Gln Glu Asp Tyr Asp
1 5 10

<210> 69
<211> 13
<212> PRT
<213> Homo sapiens

<400> 69
Leu Gly Leu Phe Asp Thr Ala Gly Gln Glu Asp Tyr Asp
1 5 10

<210> 70
<211> 13
<212> PRT
<213> Homo sapiens

<400> 70
Leu Gly Leu Trp Asp Thr Ala Gly Gln Glu Asp Tyr Asp
1 5 10

<210> 71
<211> 13
<212> PRT
<213> Homo sapiens

<400> 71
Leu Asp Ile Leu Asp Thr Ala Gly Gln Glu Glu Tyr Asp
1 5 10

<210> 72
<211> 13
<212> PRT
<213> Homo sapiens

<400> 72
Leu Asp Ile Leu Asp Thr Ser Gly Asn His Pro Phe Pro
1 5 10